BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

| Match: 1 Mismatch: 2 gap open: 5 gap extension: 2 | |
|---|---|
| x_dropoff: 0 expect: 10.000€ wordsize: 11 Filter □ View option Standard | ¥ |
| Masking character option X for protein, n for nucleotide Masking color option Black | |
| Show CDS translation Align | |

Sequence 1: gi|184371|Human hepsin mRNA, complete cds Length = 1783

Sequence 2: gi|14602458|Homo sapiens transmembrane protease, serine 2 (TMPRSS2), mRNA Length = 3226

No significant similarity was found

CPU time: 0.05 user secs.

0.03 sys. secs

0.08 total secs.

Structure

BLAST

Taxonomy

Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

| Match: 1 Mismatch: 2 gap open: 5 gap extension: 2 | |
|---|---|
| x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter \(\tau \) View option Standard | ~ |
| Masking character option X for protein, n for nucleotide Masking color option Black | |
| ☐ Show CDS translation Align | |

Sequence 1: gi|184371|Human hepsin mRNA, complete cds Length = 1783

Sequence 2: gi|13173470|Homo sapiens transmembrane protease, serine 3 (TMPRSS3), transcript variant A, mRNA >gi|12246823|dbj|AB038157.1| Homo sapiens TMPRSS3a mRNA for serine protease, complete cds Length = 2468

No significant similarity was found

CPU time:

0.05 user secs.

0.04 sys. secs

0.09 total secs.

OMIM

Taxonomy

Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

| Match: 1 Mismatch: 2 gap open: 5 gap extension: 2 | |
|---|--|
| x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter \(\tau \) View option \(\text{Standard} \) | |
| Masking character option X for protein, n for nucleotide Masking color option Black | |
| Show CDS translation Align | |

Sequence 1: gi|184371|Human hepsin mRNA, complete cds Length = 1783

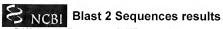
Sequence 2: gi|145701031|Homo sapiens transmembrane protease, serine 4 (TMPRSS4), transcript variant 1, mRNA Length = 3549

No significant similarity was found

CPU time: 0.05 user secs.

0.05 sys. secs

0.10 total secs.



PubMed

Entrez

BLAST

OMIM Taxon

Taxonomy

Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

| Match: 1 Mismatch: -2 gap open: 5 gap extension: 2 | |
|---|---|
| x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter □ View option Standard | V |
| Masking character option X for protein, n for nucleotide Masking color option Black | |
| Show CDS translation Align | |

Sequence 1: gi|184371|Human hepsin mRNA, complete cds Length = 1783

Sequence 2: gi|170932491|Homo sapiens transmembrane protease, serine 5 (spinesin) (TMPRSS5), mRNA Length = 2233

No significant similarity was found

CPU time:

0.04 user secs.

0.04 sys. secs

0.08 total secs.



BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

| Match: 1 Mismatch: -2 gap open: 5 gap extension: 2 | |
|---|----------|
| x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter \(\Gamma\) View option Standard | - |
| Masking character option X for protein, n for nucleotide Masking color option Black | |
| Show CDS translation Align | |

Taxonomy

Sequence 1: gi|184371|Human hepsin mRNA, complete cds Length = 1783

Sequence 2: gi|9719385|Homo sapiens enteropeptidase (PRSS7) gene, partial cds Length = 224

No significant similarity was found

CFU time: 0.05 user secs.

0.06 sys. secs

0.11 total secs.

Structure



Blast 2 Sequences results

PubMe

Entrez

DI AOT

OMIM

Taxonomy

Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTN 2.2.18 [Mar-02-2008]

| Match: 1 Mismatch: 2 gap open: 5 gap extension: 2 | |
|---|---|
| x_dropoff: 0 expect: 10.0000 wordsize: 11 Filter □ View option Standard | ~ |
| Masking character option X for protein, n for nucleotide Masking color option Black | |
| Show CDS translation Align | |

Sequence 1: gi|184371|Human hepsin mRNA, complete cds Length = 1783

Sequence 2: gi|116256362|Homo sapiens transmembrane protease, serine 13 (TMPRSS13), mRNA Length = 3391

No significant similarity was found

CPU time:

0.05 user secs.

0.04 sys. secs

0.09 total secs.